SEQUENCE LISTING	3
<pre><110> Prent, Roger</pre>	eanless of the
<pre><1100 DETECTION SYSTEMS FOR REGISTERING FROTEIN INTERACTIONS AND FUNCTIONAL RELATIONSHIPS</pre>	15 73 73 75 75 75 75 75 75 75 75 75 75 75 75 75
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<pre><140 - US 09/757,309 <141 - 2001-01-09</pre>	
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-d1605 17	TECH CENTER 1600/2900
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<pre><210 3 <211 32 <212 DNA <213 Saccharcmyces cerevisiae</pre>	
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.210	
.400 4 togagoacto cotatoagtg atagagaaaa cactocotat cagtgataga	gaaaag 56

#:210 % 5 #:211 * 56 #:213 * Excherichia coli	
6400×8 satisfying agreement to the state of the same state of	56
<pre>#210 # 6 #211 # 14 #212 DNA #213 Saccharomyces cerevisiae</pre>	
-400- e yaqoqoyata qogo	14
<pre><210 - 7 +211 - 29 <212 - DNA +313 - Homo sapiens</pre>	
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-111 - 8 -111 - 86 -1113 - DNA -113 - Homo sapiens	
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+010 × 10 <0115 33 <0105 DNA <0135 Homo sapiens	
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<pre>%210.0 11 %2110 29 %2120 DNA %2130 Baccharomyces cerevisiae</pre>	
0400% 11 qootgaatto atgtottogg totootoag	29
#210% 11 #211% 33 #211% INA	

<pre><!--213" Saccharomyces cerevisiae</pre--></pre>	
-:400 12 occgaactog agttatogaa ataacctaga agg	33
+1216 + 13 +2212 + 31 +2212 + DNA +2213 - Homo sapiens	
04000-13 godigaatto atgaaggggt ggtatcacgg a	31
+0010 + 14 +0110 + 35 +0110 + DMA +013 + Homo sapiens	
6406 14 programming agetactiga catcatiggt tittig	35
+(210) + 15 +(211 + 20) +(212) + PAT +(210) + Homo sapiens	
Asp Met Asp Trp Phe Phe Arg Phe Tyr Ala Ser Val Ser Arg Leu Phe 1 5 10 15 Arg His Leu His 23	
+210.4 10 +2115 20 +2125 PRT +213 + Homo sapiens	
Phe Trp Gln Ala Thr Leu Arg Leu Val Ser Asp Lys Leu Ile Leu Leu I 5 10 15 Tyr Fre Asp Pro 20	
#12100 17 +1.110 10 +1.110 DNA +1.110 Homo sapiens	
+400+17 ctagggccct agcatg	16